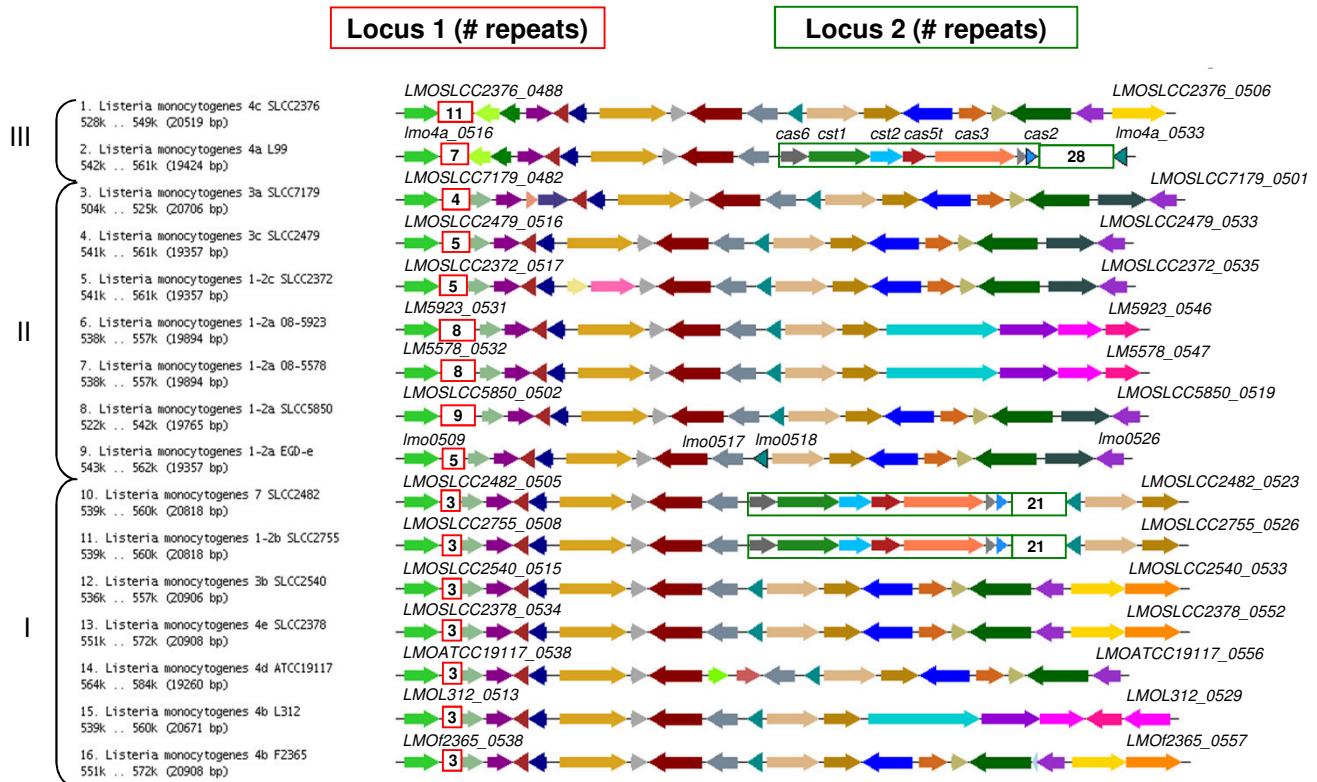


## CRISPR/Cas loci 1-3

Based on a homology cutoff >60% amino acid identity and >80% coverage. A black border denotes a deviation from the average codon usage of the chromosome. Locus 3 includes a *trans*-acting sRNA called tracrRNA that was described to compensate for a missing endoribonuclease in conjunction with host factor RNase III [Deltcheva et al., 2011, CRISPR RNA maturation by trans-encoded small RNA and host factor RNase III].

### A) Region *lmo0509-lmo0526*



### B) Region *lmo2590-lmo2614*

